## SEQUENCE LISTING

<110> Estell, David Harding, Fiona

<120> PROTEINS PRODUCING AN ALTERED IMMUNOGENIC RESPONSE AND METHODS OF MAKING AND USING THE SAME

<130> A-68893/DJB/DAV

<140> 09/500,135

<141> 2000-02-08

<150> 09/060,872

<151> 1998-04-15

<160> 236

<170> PatentIn Ver. 2.1

<210> 1

<211> 1495

<212> DNA

<213> Bacillus amyloliquefaciens

<220>

<221> mat\_peptide

<222> (417)..(1495)

<220>

<221> CDS

<222> (96)..(1244)

<220>

<221> unsure

<222> (96)..(98)

<223> The nnn at positions 96 through 98 represents gtg, which is to code for methionine.

<220>

<221> unsure

<222> (582)..(584)

<223> The nnn at positions 582 through 584 represents Xaa, which in a preferred embodiment (aat) is to code for asparagine, but which may also code for proline.

<220>

- <221> unsure
- <222> (585)..(587)
- <223> The nnn at positions 585 through 587 represents Xaa, which in a preferred embodiment (cct) is to code for proline, but which may also code for asparagine.
- <220>
- <221> unsure
- <222> (597)..(599)
- <223> The nnn at positions 597 to 599 represents Xaa, which in a preferred embodiment (aac) is to code for asparagine, but which may also code for aspartic acid.
- <220>
- <221> unsure
- <222> (678)..(680)
- <223> The nnn at positions 678 through 680 represents Xaa, which in a preferred embodiment (gca) is to code for alanine, but which may also code for serine.
- <220>
- <221> unsure
- <222> (681)..(683)
- <223> The nnn at positions 681 through 683 represents Xaa, which in a preferred embodiment (tca) is to code for serine, but which may also code for alanine.
- <220>
- <221> unsure
- <222> (708)..(710)
- <223> The nnn at positions 708 through 710 represents Xaa, which in a preferred embodiment (gct) is to code for alanine, but which may also code for aspartic acid.
- <220>
- <221> unsure
- <222> (711)..(713)
- <223> The nnn at positions 711 through 713 represents Xaa, which in a preferred embodiment (gac) is to code for aspartic acid, but which may also code for alanine.

<220>

<222>	unsur (888) The na Xaa, code serin	(8) nn a which	pos n in	a pi	refer	rred	embo	odime	ent	(act	) is	to			
<222>	unsure (891) The note three th	(8: nn a which	pos n in seri	a p	refei	rred	embo	odime	ent	(tcc	) is				
<222>	unsur (1167 The n Xaa, code for g	)(i nn a which	t pos n in gluta	sitio a pr amic	refei	rred	embo	odime	ent	(gaa	) is	to			
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ttatto	ctgca .	aatga	aaaa	aa aq	ggaga	agga	t aaa			Arg (			aaa g Lys Y		113
~ -	tc agt le Ser 00	_	_		_						_	_			161
	gc aca er Thr														209
	at att yr Ile	_				_		_	_	_	_	_	_	_	257

ttc aaa tat gta gac gca gct tca gct aca tta aac gaa aaa gct gta 353
Phe Lys Tyr Val Asp Ala Ala Ser Ala Thr Leu Asn Glu Lys Ala Val

aag aag aaa gat gtc att tct gaa aaa ggc ggg aaa gtg caa aag caa Lys Lys Asp Val Ile Ser Glu Lys Gly Gly Lys Val Gln Lys Gln



-25

-30

-35

aaa gaa ttg aaa aaa gac ccg agc gtc gct tac gtt gaa gaa gat cac 401 Lys Glu Leu Lys Lys Asp Pro Ser Val Ala Tyr Val Glu Glu Asp His -20 -15 gta gca cat gcg tac gcg cag tcc gtg cct tac ggc gta tca caa att 449 Val Ala His Ala Tyr Ala Gln Ser Val Pro Tyr Gly Val Ser Gln Ile -1 1 aaa gee eet get etg eac tet eaa gge tae aet gga tea aat gtt aaa 497 Lys Ala Pro Ala Leu His Ser Gln Gly Tyr Thr Gly Ser Asn Val Lys 2.0 gta geg gtt ate gae age ggt ate gat tet tet eat eet gat tta aag 545 Val Ala Val Ile Asp Ser Gly Ile Asp Ser Ser His Pro Asp Leu Lys 30 35 40 gta gca ggc gga gcc agc atg gtt cct tct gaa aca nnn nnn ttc caa 593 Val Ala Gly Gly Ala Ser Met Val Pro Ser Glu Thr Xaa Xaa Phe Gln 50 gac nnn aac tet cac gga aet cac gtt gee gge aca gtt geg get ett 641 Asp Xaa Asn Ser His Gly Thr His Val Ala Gly Thr Val Ala Ala Leu 60 65 70 75 aat aac tca atc ggt gta tta ggc gtt gcg cca agc nnn nnn ctt tac 689 Asn Asn Ser Ile Gly Val Leu Gly Val Ala Pro Ser Xaa Xaa Leu Tyr 80 get gta aaa gtt ete ggt nnn nnn ggt tee gge caa tae age tgg ate 737 Ala Val Lys Val Leu Gly Xaa Xaa Gly Ser Gly Gln Tyr Ser Trp Ile 100 95 att aac gga atc gag tgg gcg atc gca aac aat atg gac gtt att aac 785 Ile Asn Gly Ile Glu Trp Ala Ile Ala Asn Asn Met Asp Val Ile Asn 110 115 atg age ete gge gga eet tet ggt tet get tta aaa geg gea gtt 833 Met Ser Leu Gly Gly Pro Ser Gly Ser Ala Ala Leu Lys Ala Ala Val 125 130 135 gat aaa gee gtt gea tee gge gte gta gte gtt geg gea gee ggt aae Asp Lys Ala Val Ala Ser Gly Val Val Val Ala Ala Ala Gly Asn 140 145 150 155

gaa gge nnn nnn gge age tea age aca gtg gge tae eet ggt aaa tae

Glu Gly Xaa Xaa Gly Ser Ser Thr Val Gly Tyr Pro Gly Lys Tyr

929



160 165 170

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Ser	Ile	Gln	Ser	Thr	Leu	Pro	Gly	Asn	Lys	Tyr	Gly	Ala	Tyr	Asn	Gly	
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	Ser	Met	Ala	Ser		His	Val	Ala	GLY		Ala	Ala	Leu	Ile		
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	_				5 0					-	_	_	_	tta		1103
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aac	acc	act	aca	aaa	ctt	aat	gat	tct	ttc	tac	tat	aga	aaa	999	cta	1217
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11011			255	LID	Lou		1.00	260		- 1 -	-1-		265	O = 1		
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		270					275									
gccc	ccgc	egg t	tttt	ttat	t tt	tatt	cata	c cg	catgt	tca	atco	ggat	cca	taato	cgacgg	1324
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ggcd	caagt	cc t	gaaa	acgto	ct ca	aatco	gccg	c tto	cccg	gttt	ccgg	gtcag	gct	caato	gccgta	1444
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<211> 382

<212> PRT

<213> Bacillus amyloliquefaciens

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Ile Phe Thr Met Ala Phe Gly Ser Thr Ser Ser Ala Gln Ala Ala Gly



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Lys	Ser	Asn	Gly	Glu	Lys	Lys	Tyr	Ile	Val	Gly	Phe	Lys	Gln	Thr	Met
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Ser	Thr	Met	Ser	Ala	Ala	Lys	Lys	Lys	Asp	Val	Ile	Ser	Glu	Lys	Gly
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Gly	Lys	Val	Gln	Lys	Gln	Phe	Lys	Tyr	Val	Asp	Ala	Ala	Ser	Ala	Thr
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Leu	Asn	Glu	Lys	Ala	Val	Lys	Glu	Leu	Lys	Lys	Asp	Pro	Ser	Val	Ala
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Tyr	Val	Glu	Glu	Asp	His	Val	Ala	His	Ala	Tyr	Ala	Gln	Ser	Val	Pro
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Tyr	Gly	Val	Ser	Gln	Ile	Lys	Ala	Pro	Ala	Leu	His	Ser	Gln	Gly	Tyr
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Thr	-	Ser	Asn	Val	Lys		Ala	Val	Ile	Asp		Gly	Ile	Asp	Ser
	130		_	-		135		~ ·	~ 7		140			_	
	Hıs	Pro	Asp	Leu	-	Val	Ala	GIY	Gly		ser	Met	val	Pro	
145	Th	V	V	Db =	150	7 ~~	Vaa	7 ~~	Cox	155	C1	The se	TT = -	77-7	160
G⊥U	1111	лаа	лаа	165	GTU	Asp	лаа	ASI	Ser	птг	стλ	1111	uis	vai 175	нта
Clar	Thr	Val	ת בות		Lou	Λcn	Λαn	Sor	Ile	Clv	Val.	T ON	Clar		Nlα
GIY	1111	Val	180	Ата	Leu	ASII	ASII	185	116	GIY	vai	Leu	190	vai	AIA
Pro	Ser	Хаа		I.eu	Tur	Δla	Val		Val	Ī.e.ii	Glv	Yaa		Glv	Ser
110	bei	195	naa	БСС	- y -	nia	200	Lys	vai	Dea	Gry	205	naa	Gry	DCI
Glv	Gln		Ser	Trp	Tle	Tle		Glv	Ile	Glu	Trp		Ile	Ala	Asn
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Val	Ala	Ala	Ala	Gly	Asn	Glu	Gly	Xaa	Xaa	Gly	Ser	Ser	Ser	Thr	Val
			260					265					270		
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305					310					315				_ =	320
туг	Gly	Ala	Tyr		Gly	Thr	Ser	Met	Ala	Ser	Pro	His	Val		Gly
	• -			325		~	_		330		_		_	335	~ 7
Ala	Ala	Ala		Ile	Leu	Ser	Lys		Pro	Asn	Trp	Thr		Thr	Gln
37-3	7	<b>a</b> -	340	т -	77 -	7	m¹-	345	m¹-	т	т -	a i	350	C -	D1-
val	arg		ъer	ьeu	хаа	ASN		ınr	Thr	гуѕ	ьeu	_	Asp	ser	rne
Т••∞	Т	355	T 1.0	C1	T GVV	т 1 .	360	₹7	C1 ~	77.	<b>π</b> Ι	365	C1 ~		
τ λ.τ.	_	σтλ	ьys	στλ	ьeu		ASN	val	Gln	AIA		нта	GIN		
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<213> Bacillus amyloliquefaciens

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Ser Gly Ile Asp Ser Ser His Pro Asp Leu Lys Val Ala Gly Gly Ala 35 40 45

Ser Met Val Pro Ser Glu Thr Asn Pro Phe Gln Asp Asn Asn Ser His
50 55 60

Gly Thr His Val Ala Gly Thr Val Ala Ala Leu Asn Asn Ser Ile Gly 65 70 75 80

Val Leu Gly Val Ala Pro Ser Ala Ser Leu Tyr Ala Val Lys Val Leu 85 90 95

Gly Ala Asp Gly Ser Gly Gln Tyr Ser Trp Ile Ile Asn Gly Ile Glu 100 105 110

Trp Ala Ile Ala Asn Asn Met Asp Val Ile Asn Met Ser Leu Gly Gly
115 120 125

Pro Ser Gly Ser Ala Ala Leu Lys Ala Ala Val Asp Lys Ala Val Ala 130 135 140

Ser Gly Val Val Val Ala Ala Ala Gly Asn Glu Gly Thr Ser Gly
145 150 155 160

Ser Ser Ser Thr Val Gly Tyr Pro Gly Lys Tyr Pro Ser Val Ile Ala 165 170 175

Val Gly Ala Val Asp Ser Ser Asn Gln Arg Ala Ser Phe Ser Ser Val 180 185 190

Gly Pro Glu Leu Asp Val Met Ala Pro Gly Val Ser Ile Gln Ser Thr 195 200 205

Leu Pro Gly Asn Lys Tyr Gly Ala Tyr Asn Gly Thr Ser Met Ala Ser
210 220



Pro His Val Ala Gly Ala Ala Ala Leu Ile Leu Ser Lys His Pro Asn 225 230 235 240

Trp Thr Asn Thr Gln Val Arg Ser Ser Leu Glu Asn Thr Thr Thr Lys
245 250 255

Leu Gly Asp Ser Phe Tyr Tyr Gly Lys Gly Leu Ile Asn Val Gln Ala 260 265 270

Ala Ala Gln 275

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<212> PRT

<213> Bacillus subtilis

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His Ser Gln Gly Tyr Thr Gly Ser Asn Val Lys Val Ala Val Ile Asp 20 25 30

Ser Gly Ile Asp Ser Ser His Pro Asp Leu Asn Val Arg Gly Gly Ala 35 40 45

Ser Phe Val Pro Ser Glu Thr Asn Pro Tyr Gln Asp Gly Ser Ser His 50 55 60

Gly Thr His Val Ala Gly Thr Ile Ala Ala Leu Asn Asn Ser Ile Gly 65 70 75 80

Val Leu Gly Val Ser Pro Ser Ala Ser Leu Tyr Ala Val Lys Val Leu 85 90 95

Asp Ser Thr Gly Ser Gly Gln Tyr Ser Trp Ile Ile Asn Gly Ile Glu 100 105 110

Trp Ala Ile Ser Asn Asn Met Asp Val Ile Asn Met Ser Leu Gly Gly
115 120 125

Pro Thr Gly Ser Thr Ala Leu Lys Thr Val Val Asp Lys Ala Val Ser 130 135 140

Ser Gly Ile Val Val Ala Ala Ala Gly Asn Glu Gly Ser Ser Gly
145 150 155 160

Ser Thr Ser Thr Val Gly Tyr Pro Ala Lys Tyr Pro Ser Thr Ile Ala 165 170 175

Val Gly Ala Val Asn Ser Ser Asn Gln Arg Ala Ser Phe Ser Ser Ala 180 185 190

Gly Ser Glu Leu Asp Val Met Ala Pro Gly Val Ser Ile Gln Ser Thr 195 200 205

Leu Pro Gly Gly Thr Tyr Gly Ala Tyr Asn Gly Thr Ser Met Ala Thr 210 215 220

Pro His Val Ala Gly Ala Ala Ala Leu Ile Leu Ser Lys His Pro Thr 225 230 235 235

Trp Thr Asn Ala Gln Val Arg Asp Arg Leu Glu Ser Thr Ala Thr Tyr 245 250 255

Leu Gly Asn Ser Phe Tyr Tyr Gly Lys Gly Leu Ile Asn Val Gln Ala 260 265 270

Ala Ala Gln 275

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Gln Ala Gln Gly Phe Lys Gly Ala Asn Val Lys Val Ala Val Leu Asp 20 25 30

Thr Gly Ile Gln Ala Ser His Pro Asp Leu Asn Val Val Gly Gly Ala
35 40 45

Ser Phe Val Ala Gly Glu Ala Tyr Asn Thr Asp Gly Asn Gly His Gly 50 55 60

Thr His Val Ala Gly Thr Val Ala Ala Leu Asp Asn Thr Thr Gly Val 65 70 75 80

Leu Gly Val Ala Pro Ser Val Ser Leu Tyr Ala Val Lys Val Leu Asn

90 95

Ser Ser Gly Ser Gly Ser Tyr Ser Gly Ile Val Ser Gly Ile Glu Trp
100 105 110

Ala Thr Thr Asn Gly Met Asp Val Ile Asn Met Ser Leu Gly Gly Ala 115 120 125

Ser Gly Ser Thr Ala Met Lys Gln Ala Val Asp Asn Ala Tyr Ala Arg 130 135 140

Gly Val Val Val Val Ala Ala Ala Gly Asn Ser Gly Asn Ser Gly Ser 145 150 155 160

Thr Asn Thr Ile Gly Tyr Pro Ala Lys Tyr Asp Ser Val Ile Ala Val 165 170 175

Gly Ala Val Asp Ser Asn Ser Asn Arg Ala Ser Phe Ser Ser Val Gly
180 185 190

Ala Glu Leu Glu Val Met Ala Pro Gly Ala Gly Val Tyr Ser Thr Tyr 195 200 205

Pro Thr Asn Thr Tyr Ala Thr Leu Asn Gly Thr Ser Met Ala Ser Pro 210 215 220

His Val Ala Gly Ala Ala Ala Leu Ile Leu Ser Lys His Pro Asn Leu 225 230 235 240

Ser Ala Ser Gln Val Arg Asn Arg Leu Ser Ser Thr Ala Thr Tyr Leu 245 250 255

Gly Ser Ser Phe Tyr Tyr Gly Lys Gly Leu Ile Asn Val Glu Ala Ala 260 265 270

Ala Gln

<210> 6

<211> 269

<212> PRT

<213> Bacillus lentus

<400> 6

Ala Gln Ser Val Pro Trp Gly Ile Ser Arg Val Gln Ala Pro Ala Ala 1 5 10 15



His Asn Arg Gly Leu Thr Gly Ser Gly Val Lys Val Ala Val Leu Asp Thr Gly Ile Ser Thr His Pro Asp Leu Asn Ile Arg Gly Gly Ala Ser Phe Val Pro Gly Glu Pro Ser Thr Gln Asp Gly Asn Gly His Gly Thr His Val Ala Gly Thr Ile Ala Ala Leu Asn Asn Ser Ile Gly Val Leu Gly Val Ala Pro Ser Ala Glu Leu Tyr Ala Val Lys Val Leu Gly Ala Ser Gly Ser Gly Ser Val Ser Ser Ile Ala Gln Gly Leu Glu Trp Ala Gly Asn Asn Gly Met His Val Ala Asn Leu Ser Leu Gly Ser Pro Ser Pro Ser Ala Thr Leu Glu Gln Ala Val Asn Ser Ala Thr Ser Arg Gly Val Leu Val Val Ala Ala Ser Gly Asn Ser Gly Ala Gly Ser Ile Ser Tyr Pro Ala Arg Tyr Ala Asn Ala Met Ala Val Gly Ala Thr Asp Gln Asn Asn Asn Arg Ala Ser Phe Ser Gln Tyr Gly Ala Gly Leu Asp Ile Val Ala Pro Gly Val Asn Val Gln Ser Thr Tyr Pro Gly Ser Thr Tyr Ala Ser Leu Asn Gly Thr Ser Met Ala Thr Pro His Val Ala Gly Ala Ala Ala Leu Val Lys Gln Lys Asn Pro Ser Trp Ser Asn Val Gln Ile 230 235 Arg Asn His Leu Lys Asn Thr Ala Thr Ser Leu Gly Ser Thr Asn Leu 

Tyr Gly Ser Gly Leu Val Asn Ala Glu Ala Ala Thr Arg \$260\$

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                                     10
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<210> 76 <211> 15 <212> PRT <213> Artificial Sequence

<223> Description of Artificial Sequence: Synthetic

Thr Tyr Pro Gly Ser Thr Tyr Ala Ser Leu Asn Gly Thr Ser Met
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Thr Ser Met Ala Thr Pro His Val Ala Gly Ala Ala Ala Leu Val





1 5 10 15

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Ala Thr Pro His Val Ala Gly Ala Ala Ala Leu Val Lys Gln Lys

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<210> 83

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Lys Gln Lys Asn Pro Ser Trp Ser Val Asn Gln Ile Arg Asn His

1 5 10 15

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<212> PRT

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Asn Pro Ser Trp Ser Asn Val Gln Ile Arg Asn His Leu Lys Asn
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Asn Leu Tyr Gly Ser Gly Leu Val Asn Ala Glu Ala Ala Thr Arg
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                                                          15
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Leu Leu Arg Ala Ile Pro Arg Gln Val Ala Gln Thr Leu Gln Ala
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<210> 105
<211> 15
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Ala Ile Pro Arg Gln Val Ala Gln Thr Leu Gln Ala Asp Val Leu
                                     10
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<211> 15
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
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Arg Gln Val Ala Gln Thr Leu Gln Ala Asp Val Leu Trp Gln Met

1 5 10 15

<210> 107

<211> 15

<212> PRT

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<223> Description of Artificial Sequence: Synthetic

<400> 107

Ala Gln Thr Leu Gln Ala Asp Val Leu Trp Gln Met Gly Tyr Thr
1 5 10 15

<210> 108

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<212> PRT

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<223> Description of Artificial Sequence: Synthetic

<400> 108

Leu Gln Ala Asp Val Leu Trp Gln Met Gly Tyr Thr Gly Ala Asn
1 5 10 15

<210> 109

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<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic

<400> 109

Asp Val Leu Trp Gln Met Gly Tyr Thr Gly Ala Asn Val Arg Val
1 5 10 15

<210> 110

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<220> <223> Description of Artificial Sequence: Synthetic <400> 110 Trp Gln Met Gly Tyr Thr Gly Ala Asn Val Arg Val Ala Val Phe 10 <210> 111 <211> 15 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic <400> 111 Gly Tyr Thr Gly Ala Asn Val Arg Val Ala Val Phe Asp Thr Gly 10 <210> 112 <211> 15 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic <400> 112 Gly Ala Asn Val Arg Val Ala Val Phe Asp Thr Gly Leu Ser Glu 10 <210> 113 <211> 15 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic <400> 113 Val Arg Val Ala Val Phe Asp Thr Gly Leu Ser Glu Lys His Pro 5 1 10 15



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<210> 114
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<223> Description of Artificial Sequence: Synthetic
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                                     10
<210> 115
<211> 15
<212> PRT
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Asp Thr Gly Leu Ser Glu Lys His Pro His Phe Lys Asn Val Lys
                5
                                     10
                                                          15
<210> 116
<211> 15
<212> PRT
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<400> 116
Leu Ser Glu Lys His Pro His Phe Lys Asn Val Lys Glu Arg Thr
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                                     10
                                                          15
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<223> Description of Artificial Sequence: Synthetic
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<400> 117





Lys His Pro His Phe Lys Asn Val Lys Glu Arg Thr Asn Trp Thr 1 5 10 15

<210> 118

<211> 15

<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic

<400> 118

His Phe Lys Asn Val Lys Glu Arg Thr Asn Trp Thr Asn Glu Arg
1 5 10 15

<210> 119

<211> 15

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<223> Description of Artificial Sequence: Synthetic

<400> 119

Asn Val Lys Glu Arg Thr Asn Trp Thr Asn Glu Arg Thr Leu Asp 1 5 10 15

<210> 120

<211> 15

<212> PRT

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<223> Description of Artificial Sequence: Synthetic

<400> 120

Glu Arg Thr Asn Trp Thr Asn Glu Arg Thr Leu Asp Asp Gly Leu

1 5 10 15

<210> 121

<211> 15

<212> PRT

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1 5 10 15

<210> 122 <211> 15 <212> PRT <213> Artificial Sequence

<220>

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Asn Glu Arg Thr Leu Asp Asp Gly Leu Gly His Gly Thr Phe Val
1 5 10 15

<210> 123
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<400> 123
Thr Leu Asp Asp Gly Leu Gly His Gly Thr Phe Val Ala Gly Val
1 5 10 15

<210> 124
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<223> Description of Artificial Sequence: Synthetic

<400> 124
Asp Gly Leu Gly His Gly Thr Phe Val Ala Gly Val Ile Ala Ser
1 5 10 15

<210> 125





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<211> 15
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Gly His Gly Thr Phe Val Ala Gly Val Ile Ala Ser Met Arg Glu
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<210 > 126
<211 > 15
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<400> 126
Thr Phe Val Ala Gly Val Ile Ala Ser Met Arg Glu Cys Gln Gly
                                      10
<210> 127
<211> 15
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
<400> 127
Ala Gly Val Ile Ala Ser Met Arg Glu Cys Gln Gly Phe Ala Pro
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                                     10
                                                          15
<210> 128
<211> 15
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<400> 128
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Ile Ala Ser Met Arg Glu Cys Gln Gly Phe Ala Pro Asp Ala Glu

1 5 10 15

<210> 129

<211> 15

<212> PRT

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<223> Description of Artificial Sequence: Synthetic

<400> 129

Met Arg Glu Cys Gln Gly Phe Ala Pro Asp Ala Glu Leu His Ile
1 5 10 15

<210> 130

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<223> Description of Artificial Sequence: Synthetic

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Cys Gln Gly Phe Ala Pro Asp Ala Glu Leu His Ile Phe Arg Val 2 5 10 15

<210> 131

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<212> PRT

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<223> Description of Artificial Sequence: Synthetic

<400> 131

Phe Ala Pro Asp Ala Glu Leu His Ile Phe Arg Val Phe Thr Asn 1 5 10 15

<210> 132

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<400> 135

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<210> 136 <211> 15





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<223> Description of Artificial Sequence: Synthetic
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                                     10
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Ser Tyr Thr Ser Trp Phe Leu Asp Ala Phe Asn Tyr Ala Ile Leu
                  5
                                     10
                                                          15
<210> 138
<211> 15
<212> PRT
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Ser Trp Phe Leu Asp Ala Phe Asn Tyr Ala Ile Leu Lys Lys Ile
                 5
                                     10
<210> 139
<211> 15
<212> PRT
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Leu Asp Ala Phe Asn Tyr Ala Ile Leu Lys Lys Ile Asp Val Leu
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                                     10
                                                          15
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                 5
                                     10
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<400> 148
Pro Phe Val Asp Lys Val Trp Glu Leu Thr Ala Asn Asn Val Ile
                 5
                                     10
<210> 149
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<400> 149
Asp Lys Val Trp Glu Leu Thr Ala Asn Asn Val Ile Met Val Ser
 1
                  5
                                     10
                                                          15
<210> 150
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                                      10
<210> 152
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<400> 152
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<210> 153
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                                      10
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Met Asp Val Ile Gly Val Gly Gly Ile Asp Phe Glu Asp Asn Ile
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Asp Phe Glu Asp Asn Ile Ala Arg Phe Ser Ser Arg Gly Met Thr
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Asp Asn Ile Ala Arg Phe Ser Ser Arg Gly Met Thr Thr Trp Glu
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54

Ala Arg Phe Ser Ser Arg Gly Met Thr Thr Trp Glu Leu Pro Gly
1 5 10 15

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1 5 10 15

<210> 167

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Gly Met Thr Trp Glu Leu Pro Gly Gly Tyr Gly Arg Met Lys
1 5 10 15

<210> 168

<211> 15

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<220>

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Thr Trp Glu Leu Pro Gly Gly Tyr Gly Arg Met Lys Pro Asp Ile
1 5 10 15

<210> 169

<211> 15

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1 5 10 15

<210> 177

<211> 15

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<210> 178

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<223> Description of Artificial Sequence: Synthetic

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Cys Arg Ala Leu Ser Gly Thr Ser Val Ala Ser Pro Val Val Ala
1 5 10 15

<210> 179

<211> 15

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic

<400> 179

Leu Ser Gly Thr Ser Val Ala Ser Pro Val Val Ala Gly Ala Val
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<210> 180

<211> 15

<212> PRT

<213> Artificial Sequence

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<210> 184 <211> 15

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                                    10
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<211> 15
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<223> Description of Artificial Sequence: Synthetic
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Arg Glu Leu Val Asn Pro Ala Ser Met Lys Gln Ala Leu Ile Ala
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<210> 195

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<210> 200
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<212> PRT
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<223> Description of Artificial Sequence: Synthetic
<400> 200
Gln Ile Leu Asn Ser Tyr Lys Pro Gln Ala Ser Leu Ser Pro Ser
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<210> 201
<211> 15
<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic
<400> 201
Asn Ser Tyr Lys Pro Gln Ala Ser Leu Ser Pro Ser Tyr Ile Asp
                5
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<210> 202
<211> 15
<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic
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Lys Pro Gln Ala Ser Leu Ser Pro Ser Tyr Ile Asp Leu Thr Glu
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<210> 203
<211> 15
<212> PRT
<213> Artificial Sequence
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<400> 203
Ala Ser Leu Ser Pro Ser Tyr Ile Asp Leu Thr Glu Cys Pro Tyr
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<210> 204
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Ser Pro Ser Tyr Ile Asp Leu Thr Glu Cys Pro Tyr Met Trp Pro
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Tyr Ile Asp Leu Thr Glu Cys Pro Tyr Met Trp Pro Tyr Cys Ser
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Tyr Pro Ser Asp Phe Glu Val Ile Gln Ile Lys Glu Lys Gln Lys Ala

105

90

110

85

100

Gly	Leu	Leu 115	Thr	Leu	Glu	Asp	His 120	Pro	Asn	Ile	Lys	Arg 125	Val	Thr	Pro
Gln	Arg 130	Lys	Val	Phe	Arg	Ser 135	Leu	Lys	Tyr	Ala	Glu 140	Ser	Asp	Pro	Thr
Val 145	Pro	Cys	Asn	Glu	Thr 150	Arg	Trp	Ser	Gln	Lys 155	Trp	Gln	Ser	Ser	Arg 160
Pro	Leu	Arg	Arg	Ala 165	Ser	Leu	Ser	Leu	Gly 170	Ser	Gly	Phe	Trp	His 175	Ala
Thr	Gly	Arg	His 180	Ser	Ser	Arg	Arg	Leu 185	Leu	Arg	Ala	Ile	Pro 190	Arg	Gln
Val	Ala	Gln 195	Thr	Leu	Gln	Ala	Asp 200	Val	Leu	Trp	Gln	Met 205	Gly	Tyr	Thr
Gly	Ala 210	Asn	Val	Arg	Val	Ala 215	Val	Phe	Asp	Thr	Gly 220	Leu	Ser	Glu	Lys
His 225	Pro	His	Phe	Lys	Asn 230	Val	Lys	Glu	Arg	Thr 235	Asn	Trp	Thr	Asn	Glu 240
Arg	Thr	Leu	Asp	Asp 245	Gly	Leu	Gly	His	Gly 250	Thr	Phe	Val	Ala	Gly 255	Val
Ile	Ala	Ser	Met 260	Arg	Glu	Cys	Gln	Gly 265	Phe	Ala	Pro	Asp	Ala 270	Glu	Leu
His	Ile	Phe 275	Arg	Val	Phe	Thr	Asn 280	Asn	Gln	Val	Ser	Tyr 285	Thr	Ser	Trp
Phe	Leu 290	Asp	Ala	Phe	Asn	Tyr 295	Ala	Ile	Leu	Lys	Lys	Ile	Asp	Val	Leu
Asn 305	Leu	Ser	Ile	Gly	Gly 310	Pro	Asp	Phe	Met	Asp 315	His	Pro	Phe	Val	Asp 320
Lys	Val	Trp	Glu	Leu 325	Thr	Ala	Asn	Asn	Val 330	Ile	Met	Val	Ser	Ala 335	Ile
Gly	Asn	Asp	Gly 340	Pro	Leu	Tyr	Gly	Thr 345	Leu	Asn	Asn	Pro	Ala 350	Asp	Gln
Met	Asp	Val 355	Ile	Gly	Val	Gly	Gly 360	Ile	Asp	Phe	Glu	Asp 365	Asn	Ile	Ala

Arg Phe Ser Ser Arg Gly Met Thr Trp Glu Leu Pro Gly Gly Tyr Gly Arg Met Lys Pro Asp Ile Val Thr Tyr Gly Ala Gly Val Arg Gly Ser Gly Val Lys Gly Gly Cys Arg Ala Leu Ser Gly Thr Ser Val Ala Ser Pro Val Val Ala Gly Ala Val Thr Leu Leu Val Ser Thr Val Gln Lys Arg Glu Leu Val Asn Pro Ala Ser Met Lys Gln Ala Leu Ile Ala Ser Ala Arg Arg Leu Pro Gly Val Asn Met Phe Glu Gln Gly His Gly Lys Leu Asp Leu Leu Arg Ala Tyr Gln Ile Leu Asn Ser Tyr Lys Pro Gln Ala Ser Leu Ser Pro Ser Tyr Ile Asp Leu Thr Glu Cys Pro Tyr Met Trp Pro Tyr Cys Ser Gln Pro Ile Tyr Tyr Gly Gly Met Pro Thr Val Val Asn Val Thr Ile Leu Asn Gly Met Gly Val Thr Gly Arg Ile Val Asp Lys Pro Asp Trp Gln Pro Tyr Leu Pro Gln Asn Gly Asp Asn Ile Glu Val Ala Phe Ser Tyr Ser Ser Val Leu Trp Pro Trp Ser Gly Tyr Leu Ala Ile Ser Ile Ser Val Thr Lys Lys Ala Ala Ser Trp Glu Gly Ile Ala Gln Gly His Val Met Ile Thr Val Ala Ser Pro Ala Glu Thr Glu Ser Lys Asn Gly Ala Glu Gln Thr Ser Thr Val Lys Leu Pro Ile Lys Val Lys Ile Ile Pro Thr Pro Pro Arg Ser Lys Arg Val Leu 

Trp Asp (	Gln Tyr	His Asr		Arg	Tyr	Pro	Pro 635	Gly	Tyr	Phe	Pro	Arg 640
Asp Asn 1	Leu Arg	Met Lys	Asn	Asp	Pro	Leu 650	Asp	Trp	Asn	Gly	Asp 655	His
Ile His '	Thr Asn 660	Phe Arg	Asp	Met	Tyr 665	Gln	His	Leu	Arg	Ser 670	Met	Gly
Tyr Phe	Val Glu 675	Val Leu	Gly	Ala 680	Pro	Phe	Thr	Cys	Phe 685	Asp	Ala	Ser
Gln Tyr (	Gly Thr	Leu Leu	Met 695	Val	Asp	Ser	Glu	Glu 700	Glu	Tyr	Phe	Pro
Glu Glu :	Ile Ala	Lys Let	•	Arg	Asp	Val	Asp 715	Asn	Gly	Leu	Ser	Leu 720
Val Ile :	Phe Ser	Asp Trp	Tyr	Asn	Thr	Ser 730	Val	Met	Arg	Lys	Val 735	Lys
Phe Tyr	Asp Glu 740	Asn Thr	Arg	Gln	Trp 745	Trp	Met	Pro	Asp	Thr 750	Gly	Gly
Ala Asn	Ile Pro 755	Ala Leu	Asn	Glu 760	Leu	Leu	Ser	Val	Trp 765	Asn	Met	Gly
Phe Ser 7770	Asp Gly	Leu Tyr	Glu 775	Gly	Glu	Phe	Thr	Leu 780	Ala	Asn	His	Asp
Met Tyr 7	Tyr Ala	Ser Gly		Ser	Ile	Ala	Lys 795	Phe	Pro	Glu	Asp	Gly 800
Val Val :	Ile Thr	Gln Thr	Phe	Lys	Asp	Gln 810	Gly	Leu	Glu	Val	Leu 815	Lys
Gln Glu	Thr Ala 820	Val Val	Glu	Asn	Val 825	Pro	Ile	Leu	Gly	Leu 830	Tyr	Gln
Ile Pro A	Ala Glu 835	Gly Gly	Gly	Arg 840	Ile	Val	Leu	Tyr	Gly 845	Asp	Ser	Asn
Cys Leu A	Asp Asp	Ser His	Arg 855	Gln	Lys	Asp	Cys	Phe 860	Trp	Leu	Leu	Asp

His Ser Gly Asn Arg Gln Arg Pro Pro Ser Gly Ala Gly Ser Val Thr 885 890 Pro Glu Arg Met Glu Gly Asn His Leu His Arg Tyr Ser Lys Val Leu 900 905 Glu Ala His Leu Gly Asp Pro Lys Pro Arg Pro Leu Pro Ala Cys Pro 920 Arg Leu Ser Trp Ala Lys Pro Gln Pro Leu Asn Glu Thr Ala Pro Ser 930 935 940 Asn Leu Trp Lys His Gln Lys Leu Leu Ser Ile Asp Leu Asp Lys Val 950 955 Val Leu Pro Asn Phe Arg Ser Asn Arg Pro Gln Val Arg Pro Leu Ser 965 970 Pro Gly Glu Ser Gly Ala Trp Asp Ile Pro Gly Gly Ile Met Pro Gly 985 Arg Tyr Asn Gln Glu Val Gly Gln Thr Ile Pro Val Phe Ala Phe Leu 995 1000 1005 Gly Ala Met Val Val Leu Ala Phe Phe Val Val Gln Ile Asn Lys Ala 1010 1015 1020 Lys Ser Arg Pro Lys Arg Arg Lys Pro Arg Val Lys Arg Pro Gln Leu 1025 1030 1035 1040 Met Gln Gln Val His Pro Pro Lys Thr Pro Ser Val

<210> 209

<211> 280

<212> PRT

<213> Homo sapiens

1045

<400> 209

Arg Ala Ile Pro Arg Gln Val Ala Gln Thr Leu Gln Ala Asp Val Leu

1 5 10 15

1050

Trp Gln Met Gly Tyr Thr Gly Ala Asn Val Arg Val Ala Val Phe Asp
20 25 30

Thr Gly Leu Ser Glu Lys His Pro His Phe Lys Asn Val Lys Glu Arg

Thr	Asn 50	Trp	Thr	Asn	Glu	Arg 55	Thr	Leu	Asp	Asp	Gly 60	Leu	Gly	His	Gly
Thr 65	Phe	Val	Ala	Gly	Val 70	Ile	Ala	Ser	Met	Arg 75	Glu	Cys	Gln	Gly	Phe 80
Ala	Pro	Asp	Ala	Glu 85	Leu	His	Ile	Phe	Arg 90	Val	Phe	Thr	Asn	Asn 95	Gln
Val	Ser	Tyr	Thr 100	Ser	Trp	Phe	Leu	Asp 105	Ala	Phe	Asn	Tyr	Ala 110	Ile	Leu
Lys	Lys	Ile 115	Asp	Val	Leu	Asn	Leu 120	Ser	Ile	Gly	Gly	Pro 125	Asp	Phe	Met
Asp	His	Pro	Phe	Val	Asp	Lys 135	Val	Trp	Glu	Leu	Thr 140	Ala	Asn	Asn	Val
Ile 145	Met	Val	Ser	Ala	Ile 150	Gly	Asn	Asp	Gly	Pro 155	Leu	Tyr	Gly	Thr	Leu 160
Asn	Asn	Pro	Ala	Asp 165	Gln	Met	Asp	Val	Ile 170	Gly	Val	Gly	Gly	Ile 175	Asp
Phe	Glu	Asp	Asn 180	Ile	Ala	Arg	Phe	Ser 185	Ser	Arg	Gly	Met	Thr 190	Thr	Trp
Glu	Leu	Pro 195	Gly	Gly	Tyr	Gly	Arg 200	Met	Lys	Pro	Asp	Ile 205	Val	Thr	Tyr
Gly	Ala 210	Gly	Val	Arg	Gly	Ser 215	Gly	Val	Lys	Gly	Gly 220	Cys	Arg	Ala	Leu
Ser 225	Gly	Thr	Ser	Val	Ala 230	Ser	Pro	Val	Val	Ala 235	Gly	Ala	Val	Thr	Leu 240
Leu	Val	Ser	Thr	Val 245	Gln	Lys	Arg	Glu	Leu 250	Val	Asn	Pro	Ala	Ser 255	Met
Lys	Gln	Ala	Leu 260	Ile	Ala	Ser	Ala	Arg 265	Arg	Leu	Pro	Gly	Val 270	Asn	Met
Phe	Glu	Gln 275	Gly	His	Gly	Lys	Leu 280								

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Ala Ser Ile Ser Tyr Pro Ala Arg Tyr Ala Asn Ala Met Ala Val
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<210> 212
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<400> 212
Gly Ala Ile Ser Tyr Pro Ala Arg Tyr Ala Asn Ala Met Ala Val
                 5
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<213> Artificial Sequence

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Val Phe Ser Cys Asn Ala Asn Phe Gln Arg Ile Thr Asp Phe Asp Ala

50 55 60

Lys Ser Gly Cys Glu Pro Gly Gly Val Ala Tyr Ser Cys Ala Asp Gln 65 70 75 80

Thr Pro Trp Ala Val Asn Asp Asp Phe Ala Leu Gly Phe Ala Ala Thr 85 90 95

Ser Ile Ala Gly Ser Asn Glu Ala Gly Trp Cys Cys Ala Cys Tyr Glu 100 105 110

Leu Thr Phe Thr Ser Gly Pro Val Ala Gly Lys Lys Met Val Val Gln
115 120 125

Ser Thr Ser Thr Gly Gly Asp Leu Gly Ser Asn His Phe Asp Leu Asn 130 135 140

Gly Gly Leu Pro Gly Gln Arg Tyr Gly Gly Ile Ser Ser Arg Asn Glu 165 170 175

Cys Asp Arg Phe Pro Asp Ala Leu Lys Pro Gly Cys Tyr Trp Arg Phe 180 185 190

Asp Trp Phe Lys Asn Ala Asp Asn Pro Ser Phe Ser Phe Arg Gln Val

Gln Cys Pro Ala Glu Leu Val Ala Arg Thr Gly Cys Arg Arg Asn Asp 210 215 220

Asp Gly Asn Phe Pro Ala Val Gln Ile Pro Ser Ser Ser Thr Ser Ser 225 230 235 240

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Ser Ser Pro Pro Val Gln Pro Thr Thr Pro Ser Gly Cys Thr Ala Glu 260 265 270

Arg Trp Ala Gln 275

<210> 225

<211> 18

<212> PRT

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Gly Asn Leu Asn Phe Asp Leu Lys Glu Ile Asn Asp Ile Cys Ser Gly

115 120 125

Cys Arg Gly His Asp Gly Phe Thr Ser Ser Trp Arg Ser Val Ala Asp 130 135 140

Arg Val Val Phe Thr Gly His Ser Leu Gly Gly Ala Leu Ala Thr Val 165 170 175

Ala Gly Ala Asp Leu Arg Gly Asn Gly Tyr Asp Ile Asp Val Phe Ser 180 185 190

Tyr Gly Ala Pro Arg Val Gly Asn Arg Ala Phe Ala Glu Phe Leu Thr
195 200 205

Val Gln Thr Gly Gly Thr Leu Tyr Arg Ile Thr His Thr Asn Asp Ile 210 215 220

Val Pro Arg Leu Pro Pro Arg Glu Phe Gly Tyr Ser His Ser Ser Pro 225 230 235 240

Glu Tyr Trp Ile Lys Ser Gly Thr Leu Val Pro Val Thr Arg Asn Asp 245 250 255

Ile Val Lys Ile Glu Gly Ile Asp Ala Thr Gly Gly Asn Asn Gln Pro
260 265 270

Asn Ile Pro Asp Ile Pro Ala His Leu Trp Tyr Phe Gly Leu Ile Gly
275 280 285

Thr Cys Leu 290

<210> 228

<211> 15

<212> PRT

<213> Streptomyces plicatus

<400> 228

Ile Lys Val Leu Leu Ser Val Leu Gly Asn His Gln Gly Ala Gly
1 5 10 15

<210> 229

<211> 313

_				_
< 27	12	>	ΡR	Ή.

<213> Streptomyces plicatus

<400> 22	29
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- Met Phe Thr Pro Val Arg Arg Arg Val Arg Thr Ala Ala Leu Ala Leu

  1 5 10 15
- Ser Ala Ala Ala Leu Val Leu Gly Ser Thr Ala Ala Ser Gly Ala 20 25 30
- Ser Ala Thr Pro Ser Pro Ala Pro Ala Pro Ala Pro Ala Pro Val Lys 35 40 45
- Gln Gly Pro Thr Ser Val Ala Tyr Val Glu Val Asn Asn Asn Ser Met 50 55 60
- Leu Asn Val Gly Lys Tyr Thr Leu Ala Asp Gly Gly Asn Ala Phe
  65 70 75 80
- Asp Val Ala Val Ile Phe Ala Ala Asn Ile Asn Tyr Asp Thr Gly Thr
  85 90 95
- Lys Thr Ala Tyr Leu His Phe Asn Glu Asn Val Gln Arg Val Leu Asp 100 105 110
- Asn Ala Val Thr Gln Ile Arg Pro Leu Gln Gln Gln Gly Ile Lys Val
- Leu Leu Ser Val Leu Gly Asn His Gln Gly Ala Gly Phe Ala Asn Phe 130 135 140
- Val Ala Lys Tyr Gly Leu Asp Gly Val Asp Phe Asp Asp Glu Tyr Ala 165 170 175
- Glu Tyr Gly Asn Asn Gly Thr Ala Gln Pro Asn Asp Ser Ser Phe Val 180 185 190
- His Leu Val Thr Ala Leu Arg Ala Asn Met Pro Asp Lys Ile Ile Ser 195 200 205
- Leu Tyr Asn Ile Gly Pro Ala Ala Ser Arg Leu Ser Tyr Gly Gly Val 210 215 220
- Asp Val Ser Asp Lys Phe Asp Tyr Ala Trp Asn Pro Tyr Tyr Gly Thr 225 230 235 240

Trp Gln Val Pro Gly Ile Ala Leu Pro Lys Ala Gln Leu Ser Pro Ala 245 250 255

Ala Val Glu Ile Gly Arg Thr Ser Arg Ser Thr Val Ala Asp Leu Ala 260 265 270

Arg Arg Thr Val Asp Glu Gly Tyr Gly Val Tyr Leu Thr Tyr Asn Leu 275 280 285

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Tyr Gly Ser Glu Ala Val Arg Thr Pro 305 310

<210> 230

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<213> Bacillus amyloliquefaciens

<400> 230

Gly Thr Val Ala Ala Leu Asn Asn Ser Ile Gly Val Leu Gly Val
1 5 10 15

<210> 231

<211> 15

<212> PRT

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<400> 231

Asn Gly Ile Glu Trp Ala Ile Ala Asn Asn Met Asp Val Ile Asn 1 5 10 15

<210> 232

<211> 15

<212> PRT

<213> Bacillus lentus

<400> 232

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1 5 10 15

<210> 233

<211> 15 <212> PRT <213> Bacillus lentus <400> 233 Ser Ala Glu Leu Tyr Ala Val Lys Val Leu Gly Ala Ser Gly Ser 5 10 <210> 234 <211> 17 <212> PRT <213> Bacillus lentus <400> 234 Gly Ser Ile Ser Tyr Pro Ala Arg Tyr Ala Asn Ala Met Ala Val Gly 10 5 Ala <210> 235 <211> 15 <012> PRT <213> Bacillus lentus <400> 235 Gly Ala Gly Leu Asp Ile Val Ala Pro Gly Val Asn Val Gln Ser 5 10 15 <210> 236 <211> 272 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Hybrid of Bacillus lentus and Bacillus amyloliquefaciens Ala Gln Ser Val Pro Trp Gly Ile Ser Arg Val Gln Ala Pro Ala Ala 5 10 15 His Asn Arg Gly Leu Thr Gly Ser Gly Val Lys Val Ala Val Leu Asp

25

20

Phe Val Pro Gly Glu Pro Ser Thr Gln Asp Gly Asn Gly His Gly Thr His Val Ala Gly Thr Ile Ala Ala Leu Asn Asn Ser Ile Gly Val Leu Gly Val Ala Pro Ser Ala Glu Leu Tyr Ala Val Lys Val Leu Gly Ala Ser Gly Ser Gly Ser Val Ser Ser Ile Ala Gln Gly Leu Glu Trp Ala Gly Asn Asn Gly Met His Val Ile Asn Met Ser Leu Gly Gly Ser Gly Ser Ala Ala Leu Lys Ala Ala Val Asp Lys Ala Val Ala Ser Gly Val Val Val Ala Ala Ala Gly Asn Glu Gly Thr Ser Gly Ser Ser Ser Thr Val Gly Tyr Pro Gly Lys Tyr Pro Ser Val Ile Ala Val Gly Ala Val Asp Ser Ser Asn Gln Arg Ala Ser Phe Ser Ser Val Gly Pro Glu Leu Asp Val Met Ala Pro Gly Val Ser Ile Gln Ser Thr Leu Pro Gly Asn Lys Tyr Gly Ala Tyr Asn Gly Thr Ser Met Ala Ser Pro His Val Ala Gly Ala Ala Leu Ile Leu Ser Lys His Pro Asn Trp Thr Asn Thr Gln Val Arg Ser Ser Leu Glu Asn Thr Thr Lys Leu Gly Asp 

Thr Gly Ile Ser Thr His Pro Asp Leu Asn Ile Arg Gly Gly Ala Ser

Ser Phe Tyr Tyr Gly Lys Gly Leu Ile Asn Val Gln Ala Ala Gln